

1600

RAW SEQUENCE LISTING DATE: 03/12/2004 PATENT APPLICATION: US/10/086,489 TIME: 14:54:04

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\03122004\J086489.raw

```
3 <110> APPLICANT: The Trustees of Columbia University
     5 <120> TITLE OF INVENTION: METHOD TO DETECT MUTATIONS IN NUCLEIC ACID USING A
HYBRIDIZATION-LIGATION
             PROCEDURE
     6
     8 <130> FILE REFERENCE: 0575/44012-AB
     10 <140> CURRENT APPLICATION NUMBER: 10/086,489
    11 <141> CURRENT FILING DATE: 2002-02-28
    13 <150> PRIOR APPLICATION NUMBER: 09/100,707
    14 <151> PRIOR FILING DATE: 1998-06-19
    16 <150> PRIOR APPLICATION NUMBER: 08/853,000
     17 <151> PRIOR FILING DATE: 1997-05-08
    19 <150> PRIOR APPLICATION NUMBER: 08/409,644
                                                           ENTERED
    20 <151> PRIOR FILING DATE: 1995-03-24
    22 <160> NUMBER OF SEQ ID NOS: 7
     24 <170> SOFTWARE: PatentIn version 3.1
    26 <210> SEQ ID NO: 1
     27 <211> LENGTH: 24
     28 <212> TYPE: DNA
     29 <213> ORGANISM: human
     31 <400> SEQUENCE: 1
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     35 <210> SEQ ID NO: 2
    36 <211> LENGTH: 25
     37 <212> TYPE: DNA
     38 <213> ORGANISM: human
     40 <400> SEQUENCE: 2
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     41 agcccggtaa tcgcataaaa cttaa
     44 <210> SEQ ID NO: 3
     45 <211> LENGTH: 19
     46 <212> TYPE: DNA
     47 <213> ORGANISM: human
     49 <400> SEQUENCE: 3
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    54 <211> LENGTH: 21
    55 <212> TYPE: DNA
    56 <213> ORGANISM: human
    58 <400> SEQUENCE: 4
                                                                               21
    59 gttttatgcg attaccgggc c
    62 <210> SEQ ID NO: 5
    63 <211> LENGTH: 12
     64 <212> TYPE: DNA
    65 <213> ORGANISM: human
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67 <220> FEATURE:

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68 <221> NAME/KEY: misc feature
     69 <222> LOCATION: (1)..(1)
    70 <223> OTHER INFORMATION: "N" means any base
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    75 <222> LOCATION: (2)..(2)
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    80 <221> NAME/KEY: misc feature
    81 <222> LOCATION: (3)..(3)
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    86 <221> NAME/KEY: misc_feature
    87 <222> LOCATION: (10)..(10)
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    93 <222> LOCATION: (11)..(11)
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    98 <221> NAME/KEY: misc feature
     99 <222> LOCATION: (12)..(12)
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W--> 104 nnnccgggn nn
                                                                           12
    107 <210> SEQ ID NO: 6
    108 <211> LENGTH: 70
    109 <212> TYPE: DNA
    110 <213> ORGANISM: human
    112 <400> SEQUENCE: 6
    60
    115 ttaccgggcc
                                                                           70
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    119 <211> LENGTH: 68
    120 <212> TYPE: DNA
    121 <213> ORGANISM: human
    123 <220> FEATURE:
    124 <221> NAME/KEY: misc_structure
    125 <222> LOCATION: (20)..(47)
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    136 <221> NAME/KEY: misc feature
    137 <222> LOCATION: (21)..(21)
    138 <223> OTHER INFORMATION: "N" means any base
    141 <220> FEATURE:
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- 142 <221> NAME/KEY: misc_feature 143 <222> LOCATION: (22)..(22) 144 <223> OTHER INFORMATION: "N" means any base 147 <220> FEATURE: 148 <221> NAME/KEY: misc_feature 149 <222> LOCATION: (23)..(23) 150 <223> OTHER INFORMATION: "N" means any base 153 <220> FEATURE: 154 <221> NAME/KEY: misc feature 155 <222> LOCATION: (24)..(24) 156 <223> OTHER INFORMATION: "N" means any base
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- 160 <221> NAME/KEY: misc_feature
- 161 <222> LOCATION: (25)..(25)
- 162 <223> OTHER INFORMATION: "N" means any base
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- 166 <221> NAME/KEY: misc feature
- 167 <222> LOCATION: (26)..(26)
- 168 <223> OTHER INFORMATION: "N" means any base
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- 172 <221> NAME/KEY: misc feature
- 173 <222> LOCATION: (27)..(27)
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- 179 <222> LOCATION: (28)..(28)
- 180 <223> OTHER INFORMATION: "N" means any base
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- 184 <221> NAME/KEY: misc_feature
- 185 <222> LOCATION: (29)..(29)
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- 191 <222> LOCATION: (30)..(30)
- 192 <223> OTHER INFORMATION: "N" means any base
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- 196 <221> NAME/KEY: misc_feature
- 197 <222> LOCATION: (31)..(31)
- 198 <223> OTHER INFORMATION: "N" means any base
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- 203 <222> LOCATION: (32)..(32)
- 204 <223> OTHER INFORMATION: "N" means any base
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- 208 <221> NAME/KEY: misc feature
- 209 <222> LOCATION: (33)..(33)
- 210 <223> OTHER INFORMATION: "N" means any base
- 213 <220> FEATURE:
- 214 <221> NAME/KEY: misc feature

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Input Set : A:\pto.da.txt

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- 215 <222> LOCATION: (34)..(34)
- 216 <223> OTHER INFORMATION: "N" means any base
- 219 <220> FEATURE:
- 220 <221> NAME/KEY: misc feature
- 221 <222> LOCATION: (35)..(35)
- 222 <223> OTHER INFORMATION: "N" means any base
- 225 <220> FEATURE:
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- 227 <222> LOCATION: (36)..(36)
- 228 <223> OTHER INFORMATION: "N" means any base
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- 233 <222> LOCATION: (37)..(37)
- 234 <223> OTHER INFORMATION: "N" means any base
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- 239 <222> LOCATION: (38)..(38)
- 240 <223> OTHER INFORMATION: "N" means any base
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- 245 <222> LOCATION: (39)..(39)
- 246 <223> OTHER INFORMATION: "N" means any base
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- 251 <222> LOCATION: (40)..(40)
- 252 <223> OTHER INFORMATION: "N" means any base
- 255 <220> FEATURE:
- 256 <221> NAME/KEY: misc feature
- 257 <222> LOCATION: (41)..(41)
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- 262 <221> NAME/KEY: misc_feature
- 263 <222> LOCATION: (42)..(42)
- 264 <223> OTHER INFORMATION: "N" means any base
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- 268 <221> NAME/KEY: misc_feature
- 269 <222> LOCATION: (43)..(43)
- 270 <223> OTHER INFORMATION: "N" means any base
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- 274 <221> NAME/KEY: misc_feature
- 275 <222> LOCATION: (44)..(44)
- 276 <223> OTHER INFORMATION: "N" means any base
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- 280 <221> NAME/KEY: misc_feature
- 281 <222> LOCATION: (45)..(45)
- 282 <223> OTHER INFORMATION: "N" means any base
- 285 <220> FEATURE:
- 286 <221> NAME/KEY: misc_feature _
- 287 <222> LOCATION: (46)..(46)

68

RAW SEQUENCE LISTING	•	DATE:	03/12/200
PATENT APPLICATION:	US/10/086,489	TIME:	14:54:04

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288 <223	3> OTHER INFORMATION: "N" means any base
291 <220	D> FEATURE:
292 <22	<pre>NAME/KEY: misc_feature</pre>
293 <222	2> LOCATION: (47)(47)
294 <223	3> OTHER INFORMATION: "N" means any base
297 <400)> SEQUENCE: 7
W> 298 ctg	ccatctt aacaaacccn nnnnnnnnn nnnnnnnnn nnnnnnngtt ttatgcgatt 60

300 accgggcc

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/086,489

DATE: 03/12/2004 TIME: 14:54:05

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 1,2,1,10,11,12 Seq#:7; N Pos. 20,24,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39 Seq#:7; N Pos. 40,41,42,43,44,45,46,46

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

VERIFICATION SUMMARY

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L:104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0 L:298 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0